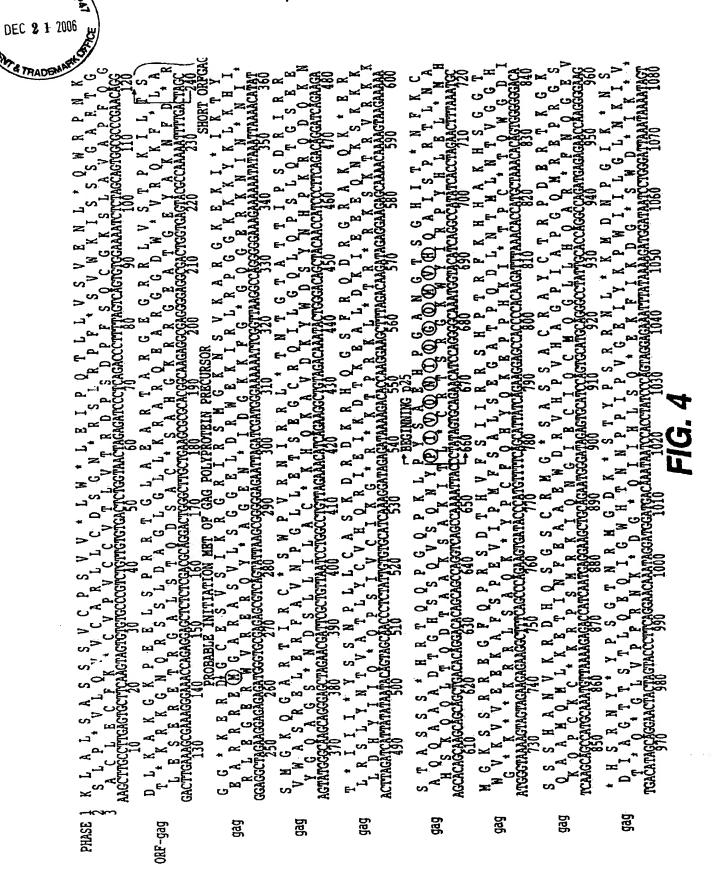


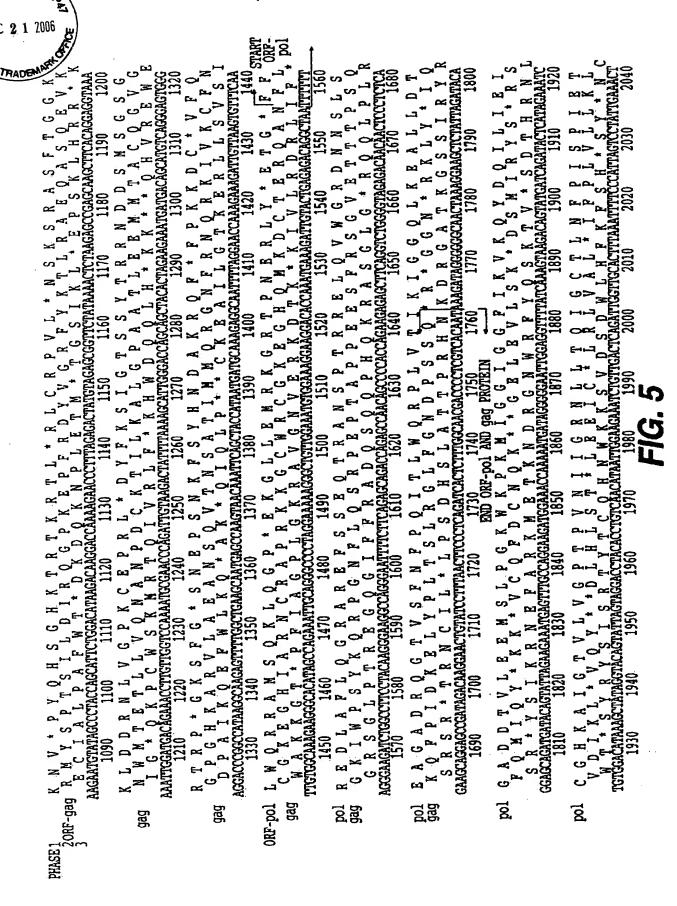




THE SECTION SE E

FIG. 3





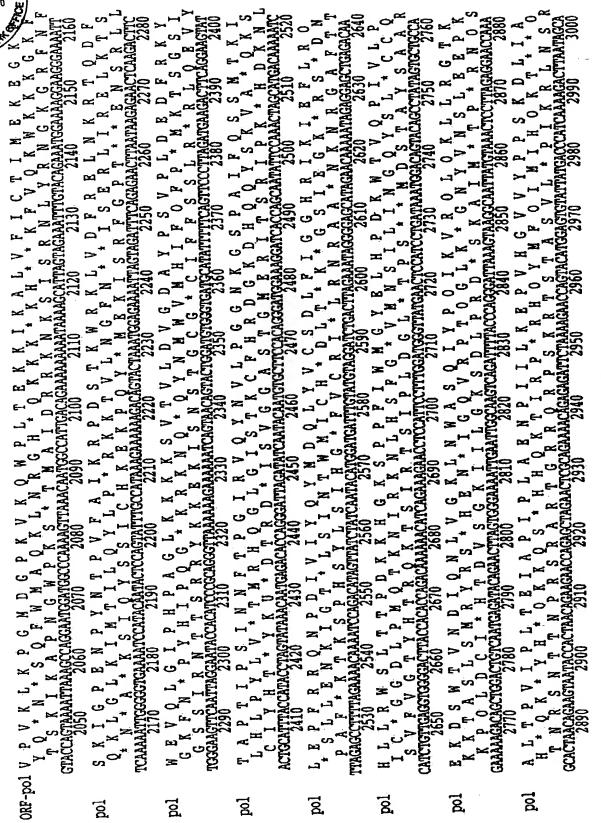
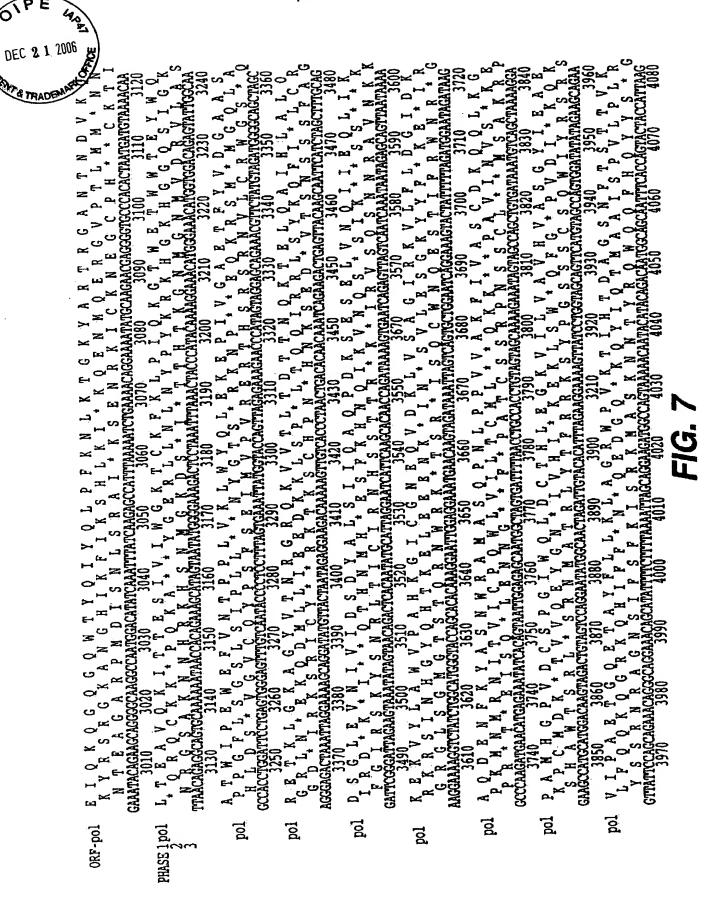
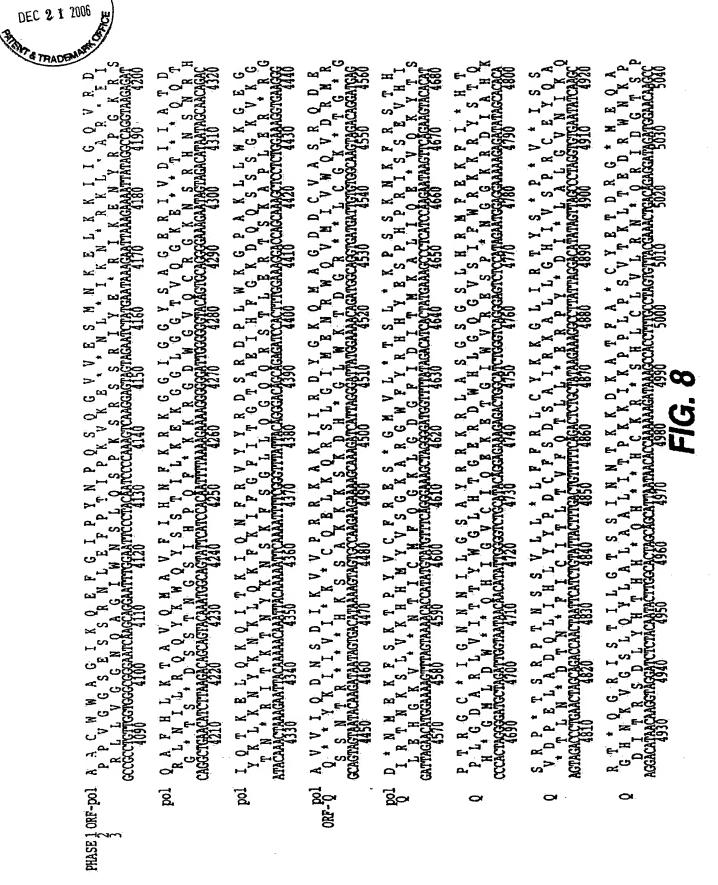
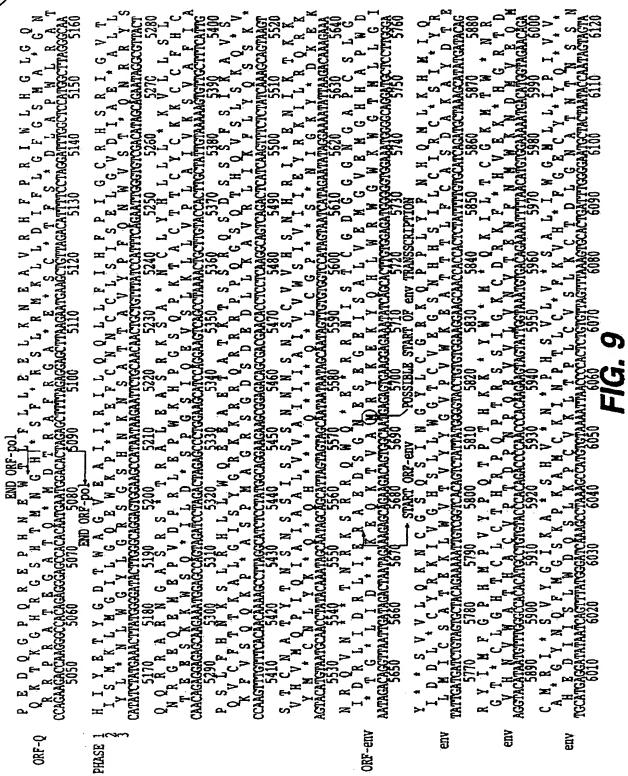


FIG. 6

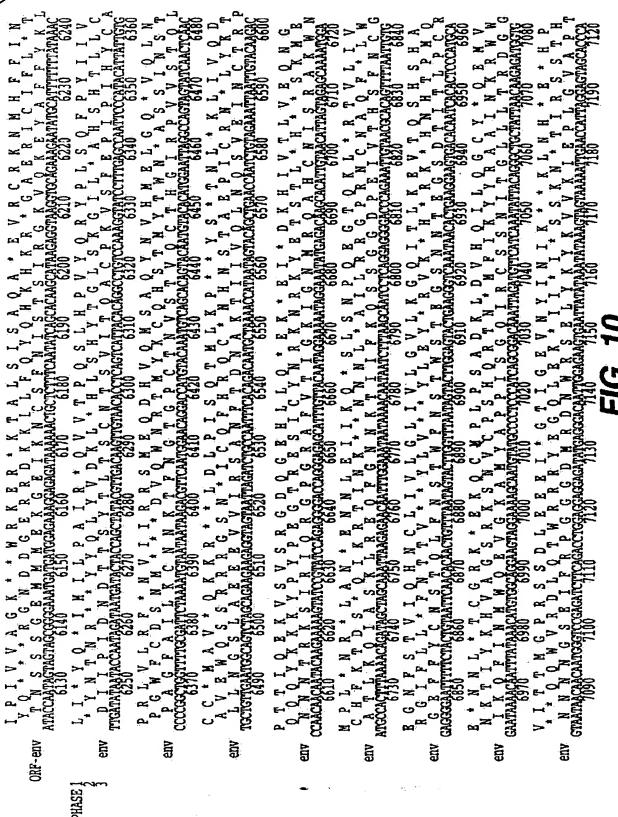


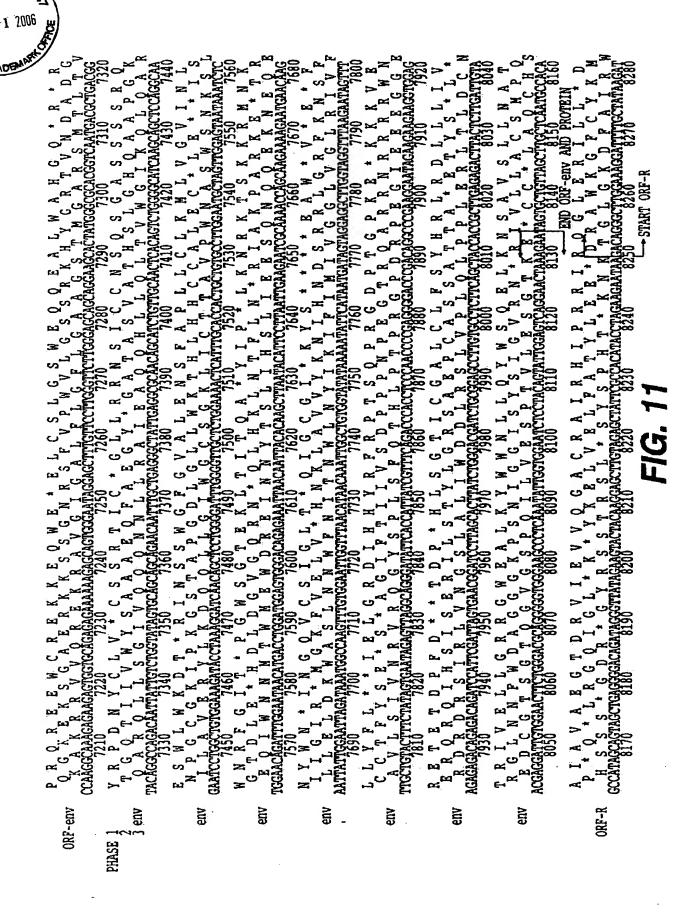












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N R G E Q E M E P V D P R L E P W K H P G S Q P K T F E S K K W S Q * I L D * S P G S I Q E V S L CAACAGAGGGAGGCAAGAAATGGAGCCAGTAGATCCTAGACTAGAGCCCTGGAAGCATCCAGGAAGTCAGCCTAA 5290 5300 5310 5320 5330 5340 5350

PSLFHNKSLRHLLWQEEAETATKTS
QVCFTTKALGISYGRKKRRQRRRPP
KFVSQQKP*ASPMAGRSGDSDEDLL
CCAAGTTTGTTTCACAACAAAAGCCTTAGGCATCTCCTATGGCAGGAAGACCGGAGACAGCGAAGACCTCCT
5410 5420 5430 5440 5450 5460 5470

S T C N A T Y T N S N S S I S S S N N N S N S C V V H V M Q P I Q I A I A A L V V A I I I A I V V W Y M C N L Y K * Q * Q H * * * Q * * * * Q * L C C AGTACATGTAATGCAACCTATACAAATAGCAATAGCAGCATTAGTAGTAGCAATAATAATAGCAATAGTTGTGTGG 5530 5540 5550 5560 5570 5580 5590

N R Q V N * * T N R K S R R Q W Q * E * R R N I S I D R L I D R L I E R A E D S G N E S E G E I S A * T G * L I D * * K E Q K T V A M R V K E K Y Q H AATAGACAGGTTAATTGATAGACTAATAGAAAGAGCAGAAGACAGTGGCAATGAGGAGAAATATCAGCA 5650 5660 5670 5680 5690 5700 5710

C M R I * S V Y G I K A * S H V * N * P H S V L V A * G Y N Q F M G S K P K A M C K I N P T L C * F H E D I I S L W D Q S L K P C V K L T P L C V S L TGCATGAGGATATAATCAGTTTATGGGATCAAAGCCTAAAGCCATGTCTAAAATTAACCCCACTCTGTGTTAGTTT 6010 6020 6030 6040 6050 6060 6070

I P I V V A G K * * W R K E R * K T A L S I S A Q Y O * * * R G N D D G E R R D K K L L F Q Y Q H K T N S S S G E M M M E K G E I K N C S F N I S T S ATACCAATAGTAGTAGCGGGGAAATGATGATGATGAGAGAAAGGAGAGATAAAAAACTGCTCTTTCAATATCAGCACAAG 6130 6140 6150 6160 6170 6180 6190

L I * Y Q * I M I L P A I R * Q V V T P Q S L H R
* Y N T N R * * Y Y Q L Y V D K L * H L S H Y T G
D I I P I D N D T T S Y T L T S C N T S V I T Q A
TTGÄTATAATACCAATAGATAÄTGATACTACCAGCTATACGTTCACAAGTTGTAACACCTCAGTCATTACACAGGG
6250 6260 6270 6280 6290 6300 6310

PRLVLRP*NVIIRRSMEQDHVQMSA

OIPE

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P G S Q P K T A C I T C Y C K K C C F H C Q E V S L K L L V P L A I V K S V A F I A CAGGAAGTCAGCCTAAAAACTGCTTGCTTGCTATTGTAAAAAGTGTTGCTTTCATTG 5370 5380 A T K T S S R Q S D S S S F S I K A V S Q R R R P P Q G S Q T H Q V S L S K Q * V S D E D L L K A V R L I K F L Y Q S S K AGCGACGAAGACCTCCTCAAGGCAGTCAGACTCATCAAGTTTCTCTATCAAAGCAGTAAGT S N S C V V H S N H R I * E N I K T K K
A I V V W S I V I I E Y R K I L R Q R K
* Q * L C G P * * S * N I G K Y * D K E K TAGCĂATAGŢŢĠŦĠŦĠĞŢĊĊĄŦAĠŦAĄŢŎĄŢAĠAÄŦĄŢĄĠĞAAÄAŦĀŢŢĄĄĠĀĊAÄĄĢĀĄĄ R R N I S T C G D G G G N G A P C S L G G E I S A L V E M G V E M G H H A P W D K E K Y Q H L W R W G W K W G T M L L G **AAGGAGAAATATCÃGCACTTGTGGAGATGGGGGTGGAAATGGGGCACCATGCTCCTTGGGA** C G P K Q P P L Y F V H Q M L K H M I Q V E G S N H H S I L C I R C * S I * Y R V W K E A T T T L F C A S D A K A Y D T E TGTGGAAGGAAGCAACCACCACTCTATTTTGTGCATCAGATGCTAAAGCATATGATACAG * Y W * M * Q K I L T C G K M T W * N R S I G K C D R K F * H V E K * H G R T D V V L V N V T E N F N M W K N D M V E Q M TAGTATTGGTAAATGTGACAGAAAATTTTAACATGTGGAAAAATGACATGGTAGAACAGA H S V L V * S A L I W G M L L I P I V V
T L C * F K V H * F G E C Y * Y O * * *
P L C V S L K C T D L G N A T N T N S S N
CACTCTGTGTTAGTTTAAAGTGCACTGATTTGGGGAATGCTACTAATACCAATAGTAGTA S I S A Q A * E V R C R K N M H F F I N O Y O H K H K R * G A E R I C I F L * T N I S I R G K V Q K E Y A F F Y K L Q S L H R P V Q R Y P L S Q F P Y I I V S H Y T G L S K G I L * A N S H T L L C S V I T Q A C P K V S F E P I P I H Y C A CAGTCATTACACĂGGCCTĞTCCAAÂGGTATCCTTTGĂGCCAATTCCCATACATTATTĞTĞ V Q M S A Q Y N V H M E L G Q * Y Q L N



- P C W F C D S K M * * * D V O W N R T M Y K C O P A G F A I L R C N N K T F N G T G P C T N V S CCCCGGCTGGTTTTGCGATCTAAAATGTAATAATAATAAGACGTTCAATGGAACAGGACCATGTACAAATGTCAGC 6370 6380 6390 6400 6410 6420 6430
- C C * M A V * Q K K R * * L D L P I S Q T M L K P A V E W Q S S R R R G S N * I C Q F H R Q C * N H L L N G S L A E E E V V I R S A N F T D N A K T TGCTGTTGAATGGCAGTCTAGCAGAGAGAGAGAGAGTAGTTAGATCTGCCAATTTCACAGACAATGCTAAAACCA 6490 6500 6510 6520 6530 6540 6550
- PTTIQEKVSVSRGDQGEHLLQ*EK*
 QQQYKKKYPYPEGTRESICYNRKNR
 NNNTTRKSIRIQRGPGRAFVTIGKI
 CCAACAACAATACAAGAAAAAGTATCCGTATCCAGAGGGGACCAGGGAGAGCATTTGTTACAATAGGAAAAATAG
 6610 6620 6630 6640 6650 6660 6670
- E G N F S T V I Q H N C L I V L G L I V L G V L K R G I F L L * F N T T V * * Y L V * * Y L E Y * R G E F F Y C N S T O L F N S T W F N S T W S T E GAGGGGAATTTTCTACTGTAATTCAACACAACTGTTTAATAGTACTTGGTTTAATAGTACTTGGAGTACTGAAG 6850 6860 6870 6880 6890 6900 6910
- E * N N L * T C G R K * E K Q C M P L P S A D K L N K T I Y K H V A G S R K S N V C P S H Q R T N * I K Q F I N M W Q E V G K A M Y A P P I S G Q I GAATAAAACAATTTATAAACATGTGGGGAAGTAGGAAAAGCAATGTATGCCCCTCCCATCAGCGGACAAATTA 6970 6980 6990 7000 7010 7020 7030
- V I T T M G P R S S D L E E E I * G T I G E V N Y
 * * Q O W V R D L Q T W R R R Y E G Q L E K * I I
 N N N N N G S E I F R P G G G D M R D N W R S E L
 GTAATAACAACAATGGTCCGAGATCTTCAGACCTGGAGGAGGAGATATGAGGGACAATTGGAGAAGTGAATTAT
 7090 7100 7110 7120 7130 7140 7150
- PRQREEWCREKKEQWE*ELCSLGSWQGKEKSGAERKKSSGNRSFVPWVLGKAKRRVVQREKRAVGIGALFLGFLCCAAGGCAAAGAGAGAAGAGAGAGAAAAAAGAGCAGTGGGAATAGGAGCTTTGTTCCTTGGGTTCTTGG7210 7220 7230 7240 7250 7260 7270
- Y R P D N Y C L V * C S S R T I C * G L L R R N S T G Q T I I V W Y S A A A E Q F A E G Y * G A T A Q A R Q L L S G I V Q Q Q N N L L R A I E A Q Q TACAGGCCAGACAATTATTGTCTGGTATAGTGCAGCAGCAGCAACAATTTGCTGAGGGCTATTGAGGCGCAACAGC 7330 7340 7350 7360 7370 7380 7390
- ESWLWKDT * RINSSWGFGVALENSF



N R T M Y K C O H S T M Y T W N * N S S I N S T T T G P C T N V S T V O C T H G I R P V V S T Q L AACAGGACCATGTACAAATGTCAGCACAGTACAATGTACACATGGAATTAGGCCAGTAGTATCAACTCAAC 6420 6430 6440 6450 6460 6470 6480

F H L L Q * E K * E I * D K H I V T L V E Q N G S I C Y N R K N R K Y E T S T K * H * * S K M E A F V T I G K I G N M R Q A H C N I S R A K W N AGCATTGTTACAATAGGAAAATAGGAAATATGAGACAAGCACATTGTAACATTAGTAGAGCAAAATGGA 6660 6670 6670 6720

I I K Q * S L S N P Q E G T Q K L * R T V L I V * N N N L * A I L R R G P R N C N A Q F * L W N K I I F K Q S S G G D P E I V T H S F N C G TAATAAACAATAATCTTTAAGCAATCCTCAGGAGGGGACCCAGAAATTGTAACGCACAGTTTTAATTGTG 6780 6790 6800 6810 6820 6830 6840

C L I V L G V L K G Q I T L K E V T Q S H S H A V * * Y L E Y * R V K * H * R K * H N H T P M Q P N S T W S T E G S N N T E G S D T I T L P C R TTTTAATAGTACTGAAGGAAGTGACACAATCACACTCCCATGCA 6900 6910 6920 6930 6940 6950 6960

M P L P S A D K L D V H Q I L Q G C Y * Q E M V C P S H Q R T N * M F I K Y Y R A A I N K R W W A P P I S G Q I R C S S N I T G L L L T R D G G TGCCCTCCCATCAGCGGACAAATTAGATGTTCATCAAATATTACAGGGCTGCTATTAACAAGAGATGGTG 7020 7030 7040 7050 7060 7070 7080

* E L C S L G S W E Q Q E A L W A H G Q * R * R R S F V P W V L G S S R K H Y G R T V N D A D G G A L F L G F L G A A G S T M G A R S M T L T V AGGAGCTTTGTTCCTTGGGAGCAGCAGCAGGAAGCACTATGGGCGCACGGTCAATGACGCTGACGG 7260 7270 7280 7290 7300 7310 7320

C * G L L R R N S I C C N S Q S G A S S S S R Q A E G Y * G A T A S V A T H S L G H Q A A P G K L R A I E A Q Q H L L Q L T V W G I K Q L Q A R GCTGAGGGCTATTGAGGGCGCAACAGCATCTGTTGCAACTCACAGTCTGGGGCATCAAGCAGCTCCAGGCAA 7380 7390 7400 7410 7420 7430 7440

G V A L E N S F A P L L C L G M L V G V I N L

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ATTO TANDONS

N P G C G K I P K G S T A P G D L G L L W K T H
I L A V E R Y L K D O O L L G I W G C S G K L I
GAATCCTGGCTGTGGAAAGATACCTAAAGGATCAACAGCTCCTGGGGATTTGGGGTTGCTCTGGAAAACTCATI W N R F G I T * P G W S G T E K L T I T Q A * Y I G T D L E * H D L D G V G Q R N * Q L H K L N T E Q I W N N N M T W M E W D R E I N N Y T S L I H TGGAACAGATTTGGAATAACATGACCTGGATGGAGTGGGACAGAGAAATTAACAATTACACAAGCTTAATACAT N Y W N * I N G Q V C S I G L T * Q I G C G I * I I I G I R * M G K F V E L V * H N K L A V V Y K L L E L D K W A S L W N W F N I T N W L W Y I K AATTATTGGAATTAGATAAATGGGCAAGTTTGTGGAATTGGTTTAACATAACAAATTGGCTGTGGTATATAAAA L L Y F L * * I E L G R D I H H Y R F R P T S Q P C C T F Y S E * S * A G I F T I I V S D P P P N A V L S I V N R V R Q G Y S P L S F Q T H L P T TTGCTGTACTTTCTATAGTGAATAGAGTTAGGCÄGGGATATTCACCATTATCGTTTCÄGACCCACCTCCCAACC R E T E T D P F D * * T D P * H L S G T I C G A L E R Q R Q I H S I S E R I L S T Y L G R S A E P R D R D R S I R L V N G S L A L I W D D L R S L AGAGAGACAGACCATTCGATTAGTGAACGGATCCTTAGCACTTATCTGGGACGATCTGCGGAGCCTT TRIVELLGRRGWEALKYWWNLLQYWRGLWNFWDAGGGKPSNIGGISYSI EDCGTSGTQGVGSPQILVESPTVL ACGAGGATTGTGGAACTTCTGGGACGCÄGGGGGTGGGAAGCCCTCÄAATATTGGTGGAATCTCCTACAGTATTC I A V A E G T D R V I E V V Q G A C R A I R H I

A Q * L R G Q I G L * K * Y K E L V E L F A T

H S S S * G D R * G Y R S S T R S L * S Y S P H GCCATAGCAGTAGCTGAGGGGACAGATAGGGTTATAGAAGTAGTACAAGGAGCTTGTAGAGCTATTCGCCACAT G W Q V V K K * C G W M A Y C K G K N E T S * A S G G K W S K S S V V G W P T V R E R M R R A E P V A S G Q K V V W L D G L L * G K E * D E L S Q GGGTGGCAAGTGGTCAAAAAGTAGTGTGGTTGGATGGCCTACTGTAAGGGAAAGAATGAGACGAGCTGAGCCAG

D G S C R S * P L F K R K G G T G R A N S L P I K

DEC 2 1 2006 M

W K T H L H H C C A L E C * L E * * I S G K L I C T T A V P W N A S W S N K S L L E * CTGGAAAACTCATTTGCACCACTGCTGTGCCTTGGAATGCTAGTTGGAGTAATAAATCTC 7520 7540 7510 7530 7550 7560) A * Y I P * L K N R K T S K K R M N K K L N T F L N * R I A K P A R K E * T R S L I H S L I E E S Q N Q Q E K N E Q E CAAGCTTAATACATTCCTTAATTGAAGAATCGCÃAAACCÃGCÃAGAAAAGAATGAACÃAG 7650 7660 C G I * K Y S * * * * E A W * V * E * F V V Y K N I H N D S R R L G R F K N S F W Y I K I F I M I V G G L V G L R I V F TGTGGTATATAAAAATATTCATAATGATAGTAGGAGGCTTGGTAGGTTTAAGAATAGTTT 7760 7790 7770 7780 T S Q P R G D P T G P K E * K K K V E P P P N P E G T R Q A R R N R R R W R H L P T P R G P D R P E G I E E E G G E CCACCTCCCAACCCCGAGGGGACCCGACAGGCCCGAAGGAATAGAAGAAGAAGAAGGTGGAG C G A L C L F S Y H R L R D L L L I V S A E P C A S S A T T A * E T Y S * L * L R S L V P L Q L P P L E R L T L D C N TCTGCGGAGCCTTGTGCCTCTTCAGCTACCACCGCTTGAGAGACTTACTCTTGATTGTA L L Q Y W S Q E L K N S A V S L L N A T S Y S I G V R N * R I V L L A C S M P Q P T V L E S G T K E * C C * L A Q C H S TCCTACAGTATTGGAGTCAGGAACTAAAGAATAGTGCTGTTAGCTTGCTCAATGCCACA 8110 . 8150 8120 8130 8140 A I R H I P R R I R Q G L E R I L L * D L F A T Y L E E * D R A W K G F C Y K M Y S P H T * K N K T G L G K D F A I R W CTATTCGCCACATACCTAGAAGAATAAGACAGGGCTTGGAAAGGATTTTGCTATAAGAT 8240 8250 8260 8390 8360 8370 8380 Q G G G G F S S H T S G T F K T N D L E E E E V G F P V T P Q V P L R P M T Y R R R R W V F Q S H L R Y L * D Q * L T GAGGAGGAGGAGGTGGGTTTCCAGTCACACCTCAGGTACCTTTAAGACCAATGACTTA 8470 8480 8490 8500 8510

L P T K T R Y P * S V D L P H T R L L

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Replacement Sheet

10	20	30	40	50	60
10				TTGTGTGACT	
		AAGTAGTGTG	100	110	120
70	80 AGACCCTTTT	AGTCAGTGTG			CCCGAACAGG
				170	180
130	140	150	160		GCTTGCTGAA
GACTTGAAAG	CGAAAGGGAA			GCAGGACTCG 230	240
190	200	210	220		TTTGACTAGC
GCGCGCACGG			CTGGTGAGTA		300
250	260	270	280	290	
	AGGAGAGA			TTAAGCGGGG	GAGAATTAGA
310	320	330	340	350	360
	AAAATTCGGT		GGCAAAGAAA		TAAAACATAT
370	380	390	400	410	420
	AGCAGGGAGC		CGCTGTTAAT	CCTGGCCTGT	TAGAAACATC
430	440	450	460	470	480
	,	TGGGACAGCT		CTTCAGACAG	GATCAGAAGA
490	500	510	520	530	540
ACTTAGATCA				GTGCATCAAA	
550	560	570	580	590	600
		,	AGAGGAAGAG		GTAAGAAAAA
610	620	630	640	650	660
		ACACAGGACA			ATTACCCTAT
670	680	690	700	710	720
a contract of the contract of			TCAGGCCATA	* 41144 - 11141-	CTTTAAATGC
730	740	750	760	770	780
			CAGCCCAGAA		TGTTTTCAGC
790	800	810	820	830	840
			AAACACCATG		TGGGGGGACA
850	860	870	880	890	900
TCAAGCAGCC		TAAAAGAGAC		GAAGCTGCAG	AATGGGATAG
910	920	930	940	950	960
			ACCAGGCCAG		CAAGGGGAAG
970	980	990	1000	1010	1020
				GGATGGATGA	
1030	1040	1050	1060	1070	1080
· ·				CTGGGATTAA	
1090	1100	1110	1120	1130	1140

FIG. 19



AAGAATGTAT	AGCCCTACCA	GCATTCTGGA	CATAAGACAA	GGACCAAAAG	AACCCTTTAG
1150	1160	1170	1180	1190	1200
AGACTATGTA	GACCGGTTCT	ATAAAACTCT	AAGAGCCGAG	CAAGCTTCAC	AGGAGGTAAA
1210	1220	1230	1240	1250	1260
AAATTGGATG	ACAGAAACCT	TGTTGGTCCA	AAATGCGAAC	CCAGATTGTA	AGACTATTTT
1270	. 1280	1290	1300	1310	1320
AAAAGCATTG	GGACCAGCAG	CTACACTAGA	AGAAATGATG	ACAGCATGTC	AGGGAGTGGG
1330	1340	1350	1360	1370	1380
AGGACCCGGC	CATAAGGCAA	GAGTTTTGGC	TGAAGCAATG	AGCCAAGTAA	CAAATTCAGC
1390	1400	1410	1420	1430	1440
TACCATAATG	ATGCAAAGAG	GCAATTTTAG	GAACCAAAGA	AAGATTGTTA	AGTGTTTCAA
1450	1460	1470	1480	1490	1500
TTGTGGCAAA	GAAGGGCACA	TAGCCAGAAA	TTGCAGGGCC	CCTAGGAAAA	AGGGCTGTTG
1510	1520	1530	1540	1550	1560
GAAATGTGGA	AAGGAAGGAC	ACCAAATGAA	AGATTGTACT	GAGAGACAGG	CTAATTTTTT
1570	1580	1590	1600	1610	1620
AGGGAAGATC	TGGCCTTCCT	ACAAGGGAAG	GCCAGGGAAT	TTTCTTCAGA	GCAGACCAGA
1630	1640	1650	1660	1670	1680
GCCAACAGCC	CCACCAGAAG		GTCTGGGGTA	GAGACAACAA	CTCCCTCTCA
1690	1700	1710	1720	1730	1740
GAAGCAGGAG	CCGATAGACA	AGGAACTGTA	TCCTTTAACT	TCCCTCAGAT	CACTCTTTGG
1750	1760	1770	1780	1790	1800
	TCGTCACAAT		GGGCAACTAA	AGGAAGCTCT	ATTAGATACA
1810	1820	1830	1840	1850	1860
	ATACAGTATT	AGAAGAAATG	AGTTTGCCAG	GAAGATGGAA	ACCAAAAATG
1870	1880	1890	1900	1910	1920
	TTGGAGGTTT	TATCAAAGTA	AGACAGTATG	ATCAGATACT	CATAGAAATC
1930	1940	1950	1960	1970	1980
	AAGCTATAGG	TACAGTATTA	GTAGGACCTA	CACCTGTCAA	CATAATTGGA
1990	2000	2010	2020	2030	2040
AGAAATCTGT	TGACTCAGAT			CCATTAGTCC	
2050	2060	2070	2080	2090	2100
	AATTAAAGCC				
2110				2150	2160
	TAAAAGCATT				
2170	2180	2190	2200	2210	2220
	GGCCTGAAAA				
2230	2240	2250	2260	2270	2280
AGTACTAAAT	GGAGAAAATT				
2290	2300	2310	2320	2330	2340
	AATTAGGAAT	The state of the s			
2350	2360	2370	2380	2390	2400

GTACTGGATG	TGGGTGATGC	ATATTTTTCA	GTTCCCTTAG	ATGAAGACTT	CAGGAAGTAT
2410	2420	2430	2440	2450	2460
ACTGCATTTA				GGATTAGATA	TCAGTACAAT
2470	2480	2490	2500	2510	2520
GTGCTTCCAC	AGGGATGGAA	AGGATCACCA	GCAATATTCC	AAAGTAGCAT	GACAAAAATC
2530	2540	2550		2570	2580
TTAGAGCCTT	TTAGAAAACA			ATCAATACAT	GGATGATTTG
2590	2600	2610	2620	2630	2640
TATGTAGGAT	CTGACTTAGA	AATAGGGCAG	CATAGAACAA	AAATAGAGGA	GCTGAGACAA
2650	2660	2670	2680	2690	2700
CATCTGTTGA	GGTGGGGACT	TACCACACCA	GACAAAAAAC	ATCAGAAAGA	ACCTCCATTC
2710	2720	2730	2740	2750	2760
CTTTGGATGG	GTTATGAACT	CCATCCTGAT	AAATGGACAG	TACAGCCTAT	AGTGCTGCCA
2770	2780	2790	2800	2810	2820
GAAAAAGACA	GCTGGACTGT	CAATGACATA	CAGAAGTTAG	TGGGAAAATT	GAATTGGGCA
2830	2840	2850	2860	2870	2880
AGTCAGATTT			CAATTATGTA	AACTCCTTAG	AGGAACCAAA
2890	2900	2910	2920	2930	2940
	AAGTAATACC				AGAAAACAGA
2950	2960	2970	2980	2990	3000
, and the second				CATCAAAAGA	
3010	3020	3030	3040	3050	3060
				TTTATCAAGA	
3070	3080	3090	3100	3110	3120
AATCTGAAAA		TGCAAGAACG			TGTAAAACAA
3130	3140	3150	3160	3170	3180
TTAACAGAGG	· ·			TAATATGGGG	
3190	3200	3210	3220	3230	3240
AAATTTAAAC				GGTGGACAGA	
3250 GCCACCTGGA	3260	3270	3280	3290	3300
3310	3320	3330	3340	CTTTAGTGAA 3350	3360
				ATGTAGATGG	
3370	3380	3390	3400	3410	3420
	· -			GAGGAAGACA	
3430	3440	3450	3460	3470	3480
				CAATTCATCT	
3490	3500		3520	3530	3540
•	· ·			ATGCATTAGG	
3550	3560	3570	3580	3590	3600
				TAATAGAGCA	
3610	3620	3630	3640	3650	3660
		2000	3320	5550	5 4 0,0

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	AAGCAAAAGG		ATGGGTACCA			
	3670	3680	3690	3700	3710	3720
	GTAGATAAAT	TAGTCAGTGC				AATAGATAAG
	3730	3740	3750	3760	3770	3780
			ATATCACAGT	•		
	3790	3800	3810	3820	3830	3840
	CTGCCACCTG	TAGTAGCAAA	AGAAATAGTA	GCCAGCTGTG	,	
	3850	3860	3870	3880	3890	3900
	GAAGCCATGC	ATGGACAAGT				
	3910	3920	3930	3940	3950	3960
			GGTAGCAGTT			
	3970	3980	3990	4000	4010	4020
	GTTATTCCAG		GCAGGAAACA			
	4030	4040	4050	4060	4070	4080
	TGGCCAGTAA	AAACAATACA	TACAGACAAT	GGCAGCAATT	TCACCAGTAC	
	4090	4100	4110	4120	4130	4140
	GCCGCCTGTT	GGTGGGCGGG	AATCAAGCAG	GAATTTGGAA	TTCCCTACAA	TCCCCAAAGT
	4150	4160	4170	4180	4190	4200
	CAAGGAGTAG	TAGAATCTAT	GAATAAAGAA	TTAAAGAAAA	TTATAGGCCA	GGTAAGAGAT
	4210	4220	4230	4240	4250	4260
	CAGGCTGAAC	ATCTTAAGAC	AGCAGTACAA		TCATCCACAA	
	4270	4280	4290	4300	4310	4320
	AAAGGGGGGA	TTGGGGGGTA	CAGTGCAGGG	GAAAGAATAG	TAGACATAAT	AGCAACAGAC
	4330	4340	4350	4360	4370	4380
			AAAACAAATT	ACAAAAATTC	AAAATTTTCG	GGTTTATTAC
	4390	4400	4410	4420	4430	4440
		GAGATCCACT				
	4450	4460	4470	4480	4490	4500
	GCAGTAGTAA					•
	4510	4520	4530	4540	4550	4560
			GATGGCAGGT			
	4570	4580	4590	4600	4610	4620
			TAGTAAAACA			
	4630	4640	4650	4660	4670	4680
			ATGAAAGCCC			
	4690	4700	4710	4720	4730	4740
	and the second s		TGGTAATAAC			
	4750	4760	4770	4780	4790	4800
			GAGTCTCCAT			
	4810	4820	4830	4840	4850	4860
			ACCAACTAAT			
	4870	4880	4890	4900	4910	4920



CTCTGCTATA	AGAAAGGCCT	TATTAGGACA	TATAGTTAGC	CCTAGGTGTG	AATATCAAGC
4930	4940	4950	4960	2970	2980
AGGACATAAC	AAGGTAGGAT	CTCTACAATA	CTTGGCACTA	GCAGCATTAA	TAACACCAAA
4990	5000	5010	5020	5030	5040
AAAGATAAAG	CCACCTTTGC	CTAGTGTTAC	GAAACTGACA	GAGGATAGAT	GGAACAAGCC
5050	5060	5070	5080	5090	5100
CCAGAAGACC	AAGGGCCACA	GAGGGAGCCA	CACAATGAAT	GGACACTAGA	GCTTTTAGAG
5110	5120	5130	5140	5150	5160
GAGCTTAAGA	ATGAAGCTGT	TAGACATTTT	CCTAGGATTT	GGCTCCATGG	CTTAGGGCAA
5170	5180	5190	5200	5210	5220
					AAGAATTCTG
5230	5240	5250	5260	5270	5280
CAACAACTGC	TGTTTATCCA	TTTCAGAATT	GGGTGTCGAC		AGGCGTTACT
5290	5300	5310	5320	5330	5340
CAACAGAGGA	GAGCAAGAAA	TGGAGCCAGT	AGATCCTAGA	CTAGAGCCCT	GGAAGCATCC
5350	5360	5370	5380	5390	5400
AGGAAGTCAG	CCTAAAACTG	CTTGTACCAC	TTGCTATTGT	AAAAAGTGTT	GCTTTCATTG
5410	5420	5430	5440	5450	5460
			CATCTCCTAT		AGCGGAGACA
5470	5480	5490	5500	5510	5520
		GCAGTCAGAC			
5530	5540	5550	5560	5570	5580
		TACAAATAGC			
5590	5600	5610	5620	5630	5640
		TAGTAATCAT			
5650	5660	5670	5680	5690	5700
AATAGACAGG		GACTAATAGA			
5710	5720	5730	5740	5750	5760
		TGGAGATGGG			
5770	5780	5790	5800	5810	5820
		ACAGAAAAAT			
5830	5840	5850	5860	5870	5880
		ACTCTATTTT			
5890	5900	5910	5920	5930	5940
		ACACATGCCT			
5950	5960	5970	5980	5990	6000
		GAAAATTTTA			
6010	6020	6030	6040	6050	6060
		TTATGGGATC			
6070	080	6090	6100	6110	6120
		TGCACTGATT			
6130	6140	6150	6160	6170	6180

7					
		GAAATGATGA			
6190	6200	6210	6220	6230	6240
	·	AGAGGTAAGG			
6250	6260	6270	6280	6290	6300
TTGATATAAT		AATGATACTA			
6310	6320	6330	6340	63,50	6360
CAGTCATTAC	ACAGGCCTGT	CCAAAGGTAT			
6370	6380		6400	6410	6420
CCCCGGCTGG		CTAAAATGTA			
6430	6440	6450	6460	6470	6480
GTACAAATGT	CAGCACAGTA	CAATGTACAC			
6490	6500	6510	6520	6530	6540
TGCTGTTGAA	TGGCAGTCTA	GCAGAAGAAG			
6550	6560	6570	6580	6590	6600
ACAATGCTAA	AACCATAATA	GTACAGCTGA	ACCAATCTGT	AGAAATTAAT	
6610	6620	6630	6640	6650	6660
CCAACAACAA	TACAAGAAAA	AGTATCCGTA	TCCAGAGGGG	ACCAGGGAGA	GCATTTGTTA
6670	6680	6690	6700	6710	6720
CAATAGGAAA	AATAGGAAAT	ATGAGACAAG	CACATTGTAA	CATTAGTAGA	
6730	6740	6750	6760	6770	6780
ATGCCACTTT	AAAACAGATA	GCTAGCAAAT	TAAGAGAACA	ATTTGGAAAT	AATAAAACAA
6790	6800	6810	6820	6830	6840
TAATCTTTAA	GCAATCCTCA	GGAGGGGACC	CAGAAATTGT	AACGCACAGT	TTTAATTGTG
.6850	6860	6870	6880	6890	6900
GAGGGGAATT	TTTCTACTGT	AATTCAACAC	AACTGTTTAA	TAGTACTTGG	
6910	6920	6930	6940	6950	6960
CTTGGAGTAC	TGAAGGGTCA	AATAACACTG	AAGGAAGTGA		
6970	6980	6990	7000	7010	7020
GAATAAAACA	ATTTATAAAC	ATGTGGCAGG	AAGTAGGAAA	AGCAATGTAT	
7030	7040	7050	7060	7070	7080
TCAGCGGACA	AATTAGATGT	TCATCAAATA	TTACAGGGCT	GCTATTAACA	AGAGATGGTG
7090	7100	7110	7120	7130	7140
GTAATAACAA	CAATGGGTCC	GAGATCTTCA	GACCTGGAGG	AGGAGATATG	
7150	7160	7170	7180	7190	7200
GGAGAAGTGA	ATTATATAAA	TATAAAGTAG	TAAAAATTGA		
7210	7220	7230	7240	7250.	7260
CCAAGGCAAA	GAGAAGAGTG	GTGCAGAGAG		AGTGGGAATA	
7270	7280	7290	7300	7310	7320
		GCAGGAAGCA		•	
7390	7340	7350	7360	7370	7380
		TCTGGTATAG			
7390	7400	7410	7420	7430	7440



	TTGAGGCGCA	ACAGCATCTG	TTGCAACTCA	CAGTCTGGGG	CATCAAGCAG	CTCCAGGCAA
	7450	7460	7470	7480	7490	7500
	GAATCCTGGC	TGTGGAAAGA	TACCTAAAGG	ATCAACAGCT	CCTGGGGATT	TGGGGTTGCT
	7510	7520	7530	7540	7550	7560
	CTGGAAAACT	CATTTGCACC	ACTGCTGTGC	CTTGGAATGC	TAGTTGGAGT	AATAAATCTC
	7570	7580	7590	7600	7610	7620
	TGGAACAGAT	TTGGAATAAC	ATGACCTGGA	TGGAGTGGGA	CAGAGAAATT	AACAATTACA
	7630	7640	7650	7660	7670	7680
	CAAGCTTAAT	ACATTCCTTA	ATTGAAGAAT	CGCAAAACCA	GCAAGAAAAG	AATGAACAAG
	7690	7700	7710	7720	7730	7740
		ATTAGATAAA	TGGGCAAGTT		GTTTAACATA	ACAAATTGGC
	7750	7760	7770	7780	7790	7800
			ATAATGATAG			
	7810	7800	7810	7820	7830	7840
			AATAGAGTTA			
	7870	7880	7890	7900	7910	7920
			GGACCCGACA		-	
	7930	7940	7950	7960	7970	7980
			ATTCGATTAG			
	7990	8000	8010	8020	8030	8040
•			CTTCAGCTAC			
	8050	8060	8070	8080	8090	8100
	ACGAGGATTG		GGGACGCAGG			
	8110	8120	8130	8140	8150	8160
			GGAACTAAAG			
	8170	8180	8190	8200	8210	8220
			GACAGATAGG			
	8230	8240	8250	8260	8270	8280
			AAGAATAAGA			
	8290	8300	8310	8320	8330	8340
			GTAGTGTGGT			8400
	8350	8360	8370	8380	8390	
			ATGGGGTGGG			8460
	8410	8420	8430	8440	8450	
	8470		CAGCAGCTAC		8510	8520
		8480	8490 TTCCAGTCAC	8500		=•
	8530	8540	8550		8570	
			GCCACTTTTT	8560		
	8590	8600	8610	8620		8640
			ATATCCTTGA	•		
	8650	8660	8670	8680	8690	8700
	0000		0010	0,000	0000	0,00

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CCCTGATTGG	CAGAACTACA	CACCAGGGCC	AGGGGTCAGA	TATCCACTGA	CCTTTGGATG
8710	8720	8730	8740	8750	8760
GTGCTACAAG	CTAGTACCAG	TTGAGCCAGA	TAAGGTAGAA	GAGGCCAATA	AAGGAGAGAA
8770	8780	8790	8800	8810	8820
CACCAGCTTG	TTACACCCTG	TGAGCCTGCA	TGGAATGGAT	GACCCTGAGA	GAGAAGTGTT
8830	8840	8850	8860	8870	8880
AGAGTGGAGG	TTTGACAGCC	GCCTAGCATT	TCATCACGTG	GCCCGAGAGC	TGCATCCGGA
8890	8900	8910	8920	8930	8940
GTACTTCAAG	AACTGCTGAC	ATCGAGCTTG	CTACAAGGGA	CTTTCCGCTG	GGGACTTTCC
8950	8960	8970	8980	8990	9000
AGGGAGGCGT	GGCCTGGGCG	GAACTGGGGA	GTGGCGAGCC	CTCAGATGCT	GCATATAAGC
9010	9020	9030	9040	9050	9060
AGCTGCTTTT	TGCCTGTACT	GGGTCTCTCT	GGTTAGACCA	GATTTGAGCC	TGGGAGCTCT
9070	9080	9090	9100	9110	9120
CTGGCTAACT	AGGGAACCCA	CTGCTTAAGC	CTCAATAAAG	CTT	

FIG. 26